

IFWO

RAW SEQUENCE LISTING

DATE: 10/05/2004

PATENT APPLICATION: US/10/771,254

1 Patin Docket Preview

8

7 <110> APPLICANT: Flores, Heather

Lin, Tanya P.

TIME: 10:33:17

Input Set : A:\P1857R1P1.txt

Output Set: N:\CRF4\10052004\J771254.raw

```
9
        Pai, Roger
10
        Shahrokh, Zahra
        Matthews, Timothy C.
13 <120> TITLE OF INVENTION: Apo-2 LIGAND/TRAIL FORMULATIONS
15 <130> FILE REFERENCE: P1857R1P1
17 <140> CURRENT APPLICATION NUMBER: US 10/771,254
18 <141> CURRENT FILING DATE: 2004-02-03
20 <150> PRIOR APPLICATION NUMBER: US 60/338,249
21 <151> PRIOR FILING DATE: 2001-11-13
23 <150> PRIOR APPLICATION NUMBER: PCT/US02/36251
24 <151> PRIOR FILING DATE: 2002-11-12
26 <160> NUMBER OF SEQ ID NOS: 2
28 <210> SEO ID NO: 1
29 <211> LENGTH: 281
30 <212> TYPE: PRT
31 <213> ORGANISM: Homo sapiens
33 <400> SEQUENCE: 1
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35
   Cys Val Leu Ile Val Ile Phe Thr Val Leu Gln Ser Leu Cys
                     20
40
   Val Ala Val Thr Tyr Val Tyr Phe Thr Asn Glu Leu Lys Gln Met
41
                     35
                                         40
43
   Gln Asp Lys Tyr Ser Lys Ser Gly Ile Ala Cys Phe Leu Lys Glu
                                         55
44
                     50
   Asp Asp Ser Tyr Trp Asp Pro Asn Asp Glu Glu Ser Met Asn Ser
46
                                         70
47
                     65
49
   Pro Cys Trp Gln Val Lys Trp Gln Leu Arg Gln Leu Val Arg Lys
                                         85
   Met Ile Leu Arg Thr Ser Glu Glu Thr Ile Ser Thr Val Gln Glu
52
                                        100
                     95
53
   Lys Gln Gln Asn Ile Ser Pro Leu Val Arg Glu Arg Gly Pro Gln
55
56
                    110
                                        115
   Arg Val Ala Ala His Ile Thr Gly Thr Arg Gly Arg Ser Asn Thr
58
59
                                        130
                    125
   Leu Ser Ser Pro Asn Ser Lys Asn Glu Lys Ala Leu Gly Arg Lys
62
                    140
                                        145
   Ile Asn Ser Trp Glu Ser Ser Arg Ser Gly His Ser Phe Leu Ser
64
65
                    155
                                        160
   Asn Leu His Leu Arg Asn Gly Glu Leu Val Ile His Glu Lys Gly
```

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```
68
                       170
                                           175
  70
      Phe Tyr Tyr Ile Tyr Ser Gln Thr Tyr Phe Arg Phe Gln Glu Glu
  71
                      185
                                           190
      Ile Lys Glu Asn Thr Lys Asn Asp Lys Gln Met Val Gln Tyr Ile
  73
  74
                      200
                                           205
  76
      Tyr Lys Tyr Thr Ser Tyr Pro Asp Pro Ile Leu Leu Met Lys Ser
  77
                      215
                                           220
                                                                225
  79
      Ala Arg Asn Ser Cys Trp Ser Lys Asp Ala Glu Tyr Gly Leu Tyr
  80
                      230
  82
      Ser Ile Tyr Gln Gly Gly Ile Phe Glu Leu Lys Glu Asn Asp Arq
  83
                      245
                                           250
                                                                255
  85
      Ile Phe Val Ser Val Thr Asn Glu His Leu Ile Asp Met Asp His
  86
  88
      Glu Ala Ser Phe Phe Gly Ala Phe Leu Val Gly
  89
                      275
  91 <210> SEQ ID NO: 2
  92 <211> LENGTH: 1042
  93 <212> TYPE: DNA
  94 <213> ORGANISM: Homo sapiens
  96. <220> FEATURE:
  97 <221> NAME/KEY: unsure
  98 <222> LOCATION: 447
  99 <223> OTHER INFORMATION: unknown base
  101 <400> SEQUENCE: 2
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       ctgcctggct gacttacagc agtcagactc tgacaggatc atggctatga 100
       tggaggtcca ggggggaccc agcctgggac agacctgcgt gctgatcgtg 150
  106
  108
       atcttcacag tgctcctgca gtctctctgt gtggctgtaa cttacgtgta 200
  110
       ctttaccaac gagetgaage agatgcagga caagtactec aaaaqtqqca 250
  112
       ttgcttgttt cttaaaagaa gatgacagtt attgggaccc caatgacqaa 300
  114
       gagagtatga acageceetg etggeaagte aagtggeaae teegteaget 350
  116
       cgttagaaag atgattttga gaacctctga ggaaaccatt tctacagttc 400
> 118
       aagaaaagca acaaaatatt tctcccctag tgagagaaag aggtccncag 450
  120
       agagtagcag ctcacataac tgggaccaga ggaagaagca acacattgtc 500
  122
       ttetecaaac tecaagaatg aaaaggetet gggeegeaaa ataaacteet 550
  124
       gggaatcatc aaggagtggg cattcattcc tgagcaactt gcacttgagg 600
  126
       aatggtgaac tggtcatcca tgaaaaaggg ttttactaca tctattccca 650
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       aacatacttt cgatttcagg aggaaataaa agaaacaca aagaacgaca 700
  130
       aacaaatggt ccaatatatt tacaaataca caagttatcc tgaccctata 750
  132
       ttgttgatga aaagtgctag aaatagttgt tggtctaaag atgcagaata 800
  134
       tggactctat tccatctatc aagggggaat atttgagctt aaggaaaatg 850
  136
       acagaatttt tgtttctgta acaaatgagc acttgataga catggaccat 900
       gaagccagtt ttttcggggc ctttttagtt ggctaactga cctggaaaga 950
       aaaagcaata acctcaaagt gactattcag ttttcaggat gatacactat 1000
  142
       gaagatgttt caaaaaatct gaccaaaaca aacaaacaga aa 1042
```

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/771,254

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 447/

VERIFICATION SUMMARY

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L:1 M:259 W: Allowed number of lines exceeded, (1) GENERAL INFORMATION: L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:400